





Libraries . Transcripts ... Library Comp : Genomic Data

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Gene Information

Gene ID: 396578 Template Count: 6

Seq Count: 557

Clone Count: 483

Top Hit ID: g180198

Hit Species:

Homo sapiens

Hit Type:

Exact

E-Value:

0.0e+00

Pct ID:

Gene Description: Human carcinoembryonic antigen mRNA.

Genomic Data

Templates

,•		Template ID	Template Description	Top Hit ID	Hit Species
Z		396578.12	Human carcinoembryonic antigen mRNA.	g180198	Homo sapien
V		396578.6	Human BGP gene for biliary glycoprotein,	g <u>29447</u>	Homo sapien
Y		396578.10	Human carcinoembryonic antigen (CEA) gen	g <u>180206</u>	Homo sapien
Z		396578.15	Human carcinoembryonic antigen mRNA (CEA	<u>g180222</u>	Homo sapien
V	•	396578.2	Human carcinoembryonic antigen mRNA.	g <u>180198</u>	Homo sapien
Y		396578.14	Human carcinoembryonic antigen (CEA) gen	g <u>180208</u>	Homo sapien

Uncheck All

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Protein Function

Function hierarchy

Adhesion and molecular recognition Adhesion

Ig superfamily

Localized and structural proteins

Membrane

Peripheral and anchored membrane proteins

Tissue Distribution

	Clone .		Abs	Pct	Pct
Tissue Category	Count	Found in	Abund	Abund	Spec
Cardiovascular System	266190	0/68	0	0.0000	0.00
Connective Tissue	144645	1/47	1	0.0007	0.64
Digestive System	501101	61/148	342	0.0682	62.17
Embryonic Structures	106713	0/21	0	0.0000	0.00
Endocrine System	225386	1/53	1	0.0004	0.36
Exocrine Glands	254635	2/64	3	0.0012	1.09
Genitalia, Female	427284	<u>8/106</u>	17	0.0040	3.65
Genitalia, Male	448207	2/114	· 5	0.0011	1.00
Germ Cells	38282	<u>0/5</u>	0	0.0000	0.00
Hemic and Immune System	680277	4/159	49	0.0072	6.56
Liver	109378	<u>1/35</u>	2	0.0018	1.64
Musculoskeletal System	159280	0/47	0	0.0000	0.00
Nervous System	955753	0/198	0	0.0000	0.00
Pancreas	110207	4/24	26	0.0236	21.51
Respiratory System	390086	4/93	6	0.0015	1.37
Sense Organs	19256	0/8	0	0.0000	0.00
Skin	72292	<u>0/15</u> .	0	0.0000	0.00
Stomatognathic System	12923	0/10	0	0.0000	0.00
Unclassified/Mixed	120926	<u>0/13</u>	0	0.0000	0.00
Urinary Tract	279062	0/64	0	0.0000	0.00
Totals	5321883	88/1292	452	0.0091	100.00

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Gene Detail



Genes Libraries Transcripts Library Comp Genomic Data Sequences

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Gene Information

1383133 Gene ID:

Template Count: 17

Seq Count: 2876 Clone Count: 279

Top Hit ID: g35740

Hit Species:

Homo sapiens

Hit Type: Exact

E-Value:

0.0e+00Pct ID: 99

Gene Description: Human mRNA for prostate specific antigen.

Genomic Data

Templates

,		Template ID	Template Description	Top Hit ID	Hit Species
Y		1383133.7	Human mRNA for prostate specific antigen	g <u>35740</u>	Homo sapien
V	•	1383133.17	Human mRNA for prostate specific antigen	g <u>35740</u>	Homo sapien
Y		1383133.8	Human mRNA for prostate specific antigen	g <u>35740</u>	Homo sapien
V		1383133.16	Human prostate specific antigen (PA) gen	g <u>189525</u>	Homo sapien
Z		1383133.13	Human mRNA for prostate specific antigen	g <u>35740</u>	Homo sapien
Z		1383133.2	Human PSA gene for prostate specific ant	g <u>35730</u>	Homo sapien
Y		1383133.9	Human prostate-specific antigen (PA) gen	g <u>341200</u>	Homo sapien
Z		1383133.11	Human mRNA for prostate specific antigen	g <u>35720</u>	Homo sapien
~		1383133.14	Human mRNA for prostate specific antigen	g <u>35740</u>	Homo sapien
Y		1383133.12	Human prostate specific antigen gene, co	<u>g190552</u>	Homo sapien
Y		1383133.15	Human prostate specific antigen gene, co	g <u>190552</u>	Homo sapien
Y		1383133.10	Human prostate specific antigen gene, co	g <u>190552</u>	Homo sapien
Y		1383133.1	Human prostate specific antigen (PA) gen	g <u>189523</u>	Homo sapien
Y		1383133.3	Human prostate specific antigen (PA) gen	g <u>189529</u>	Homo sapien
Y		1383133.4	Human mRNA for prostate specific antigen	g <u>35740</u>	Homo sapien
Y		1383133.5	Human mRNA for prostate specific antigen	g <u>35720</u>	Homo sapien
1		1383133.6	Human mRNA for prostate specific antigen	g35740	Homo sapien

Uncheck All

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Protein Function

Enzyme hierarchy

Hydrolases

Acting on peptide bonds (peptide hydrolases) Serine endopeptidases

Function hierarchy

Protein modification and maintenance
Proteases
Serine proteases
Localized and structural proteins
Secreted and extracellular
Other secreted and extracellular

Pathway hierarchy

Metabolism

Protein metabolism

Post-translation

Protein cleavage and degradation

Tissue Distribution

	Clone		Abs	Pct	Pct
Tissue Category	Count	Found in	Abund	Abund	Spec
Cardiovascular System	266190	0/68	0	0.0000	0.00
Connective Tissue	144645	1/47	4	0.0028	0.45
Digestive System	501101	6/148	6	0.0012	0.19
Embryonic Structures	106713	1/21	1	0.0009	0.14
Endocrine System	225386	<u>1/53</u>	1	0.0004	0.06
Exocrine Glands	254635	3/64	. 4	0.0016	0.26
Genitalia, Female	427284	1/106	1	0.0002	0.03
Genitalia, Male	448207	65/114	2728	0.6086	97.07
Germ Cells	38282	0/5	0	0.0000	0.00
Hemic and Immune System	680277	1/159	5	0.0007	0.11
Liver	109378	0/35	0	0.0000	0.00
Musculoskeletal System	159280	2/47	13	0.0082	1.31
Nervous System	955753	5/198	6	0.0006	0.10
Pancreas	110207	1/24	2	0.0018	0.29
Respiratory System	390086	0/93	0	0.0000	0.00
Sense Organs	19256	0/8	0	0.0000	0.00
Skin	72292	0/15	0	0.0000	0.00
Stomatognathic System	12923	0/10	0	0.0000	0.00
Unclassified/Mixed	120926	0/13	0	0.0000	0.00
Urinary Tract	279062	0/64	0	0.0000	0.00
Totals	5321883	87/1292	2771	0.0524	100.00

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